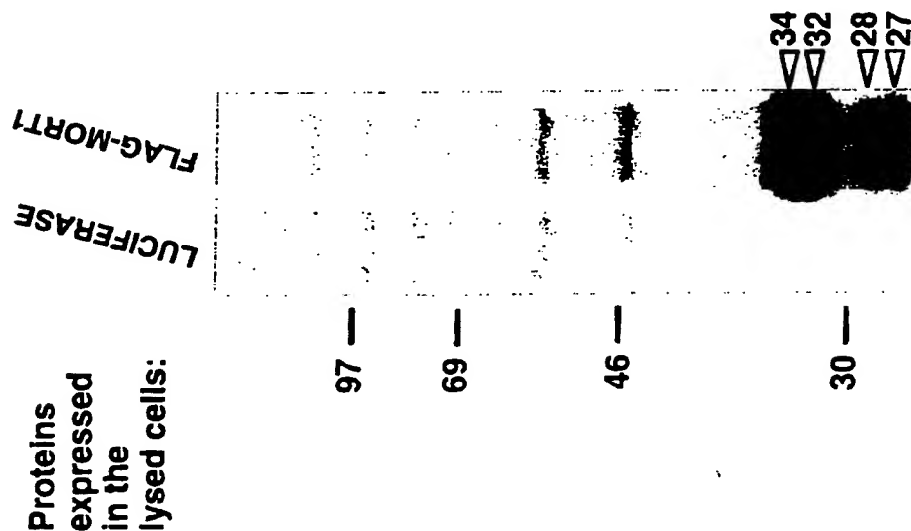


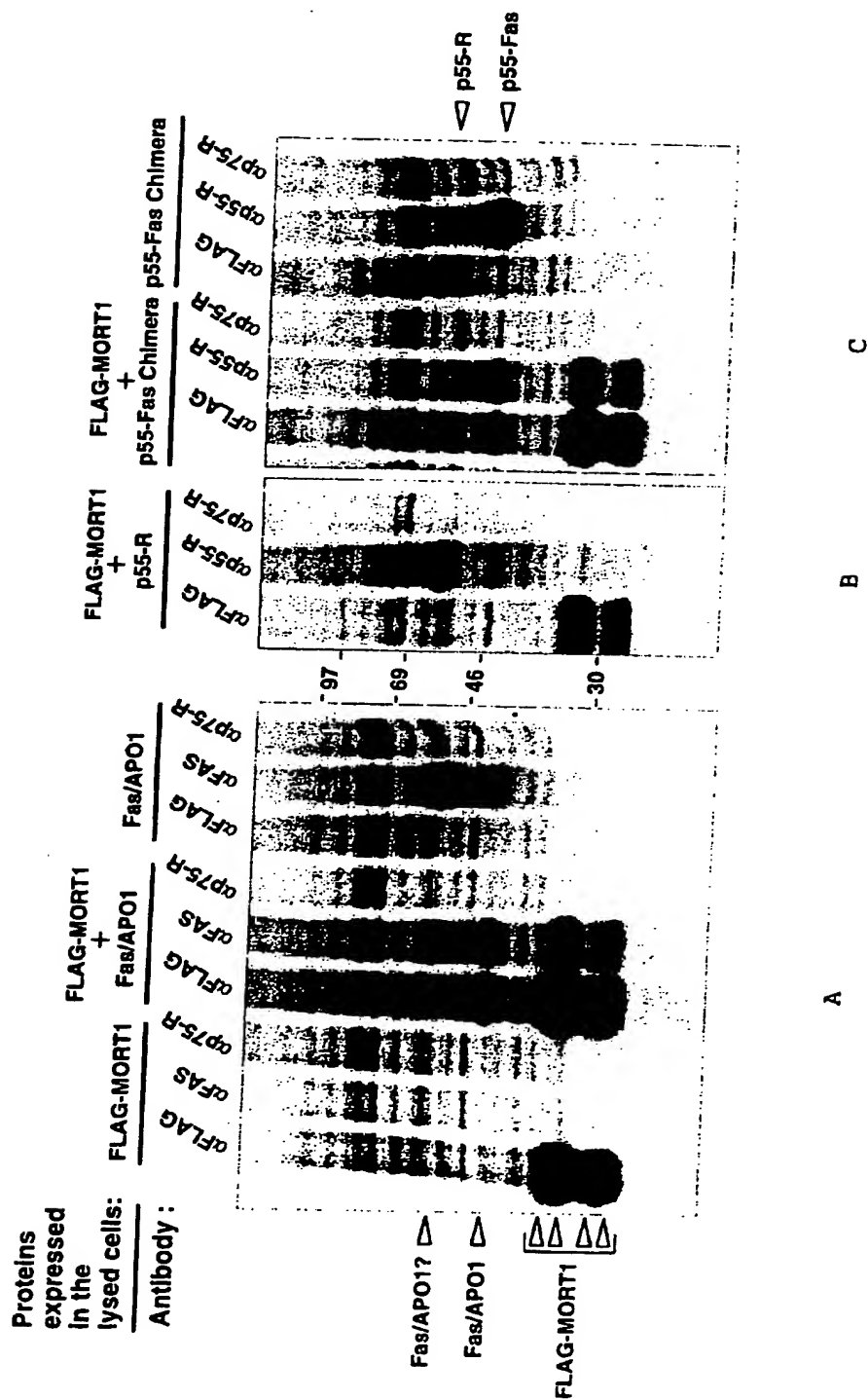
| Binding to GST fusion proteins |             |
|--------------------------------|-------------|
| GST                            | GST-mFas-IC |
| LUCIFERASE                     | LUCIFERASE  |
| FLAG-MORT1                     | FLAG-MORT1  |



A

**B**

Figure 2



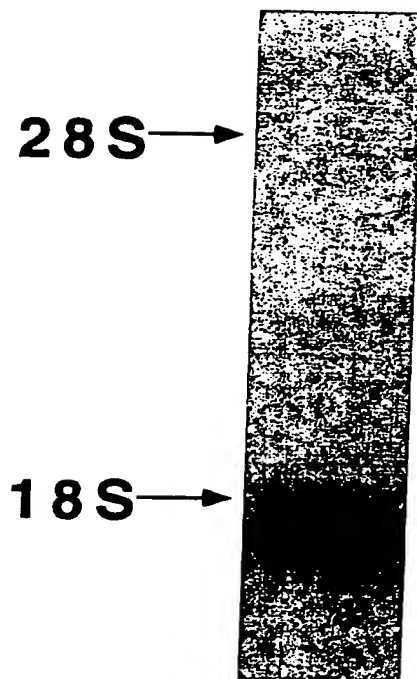


Figure 3

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1/1 31/11  
 GTG AAT CAG GCA CCG GAG TGC AGG TTC GGG GGT GGA ATC CTT GGG CCG CTG GGC AAG CCG  
 V N Q A P E C R F G G G I L G P L G K R  
 61/21 91/31  
 CGA GAC CTG GCC AGG GCC AGC GAG CCG AGG ACA GAG GCC GCG CCG AGC GCC GGG CCG CAG  
 R N L A R A S E P R T E G A R R A G P O  
 121/41 151/51  
 CCC CCG CCG CTT GCA GAC CCC GCC ATG GAC CCG TTC CTG GTG CTG CTG CAC TCG GTG TCG  
 P R P L A D P A M D P F L V L L H S V S  
 181/61 211/71  
 TCC AGC CTG TCG AGC AGC GAG CTG ACC GAG CTC AAG TTC CTA TGC CTC GCG CGC GTG GTC  
 S S L S S S E L T E L K F L C L G R V V  
 241/81 271/91  
 AAG CGC AAG CTG GAG CCG CTG CAG AGC GGC CTA GAC CTC TTC TCC ATG CTG CTG GAG CAG  
 K R K L E R V Q S G L D L F S M L L E Q  
 301/101 331/111  
 AAC GAC CTG GAG CCC GGG CAC ACC GAG CTC CTG CCG GAG CTG CTC GCC TCC CTG CCG CCG  
 N D L E P G H T E L L R E L L A S L R R  
 361/121 391/131  
 CAC GAC CTG CTG CCG CCG GTC GAC GAC TTC GAG GCG GGG GCG GCG GCC GCG CCT  
 H D L L R R V D D F E A G A A A G A A P  
 421/141 451/151  
 GGG GAA GAA GAC CTG TGT GCA GCA TTT AAC GTC ATA TGT GAT AAT GTG GGG AAA GAT TGG  
 G E E D L C A A F N V I C D N V G K D W  
 481/161 511/171  
 AGA AGG CTG GCT CGT CAG CTC AAA GTC TCA GAC ACC AAG ATC GAC AGC ATC GAG GAC AGA  
 R R L A R Q L K V S D T K I D S I E D R  
 541/181 571/191  
 TAC CCC CCG AAC CTG ACA GAG CGT GTG CCG GAG TCA CTG AGA ATC TGG AAG AAC ACA GAG  
 Y P R N L T E R V R E S L R I W K N T E  
 601/201 631/211  
 AAG GAG AAC GCA ACA GTG GCC CAC CTG GTG GGG GCT CTC AGG TCC TGC CAG ATG AAC CTG  
 K E N A T V A H L V G A L R S C Q M N L  
 661/221 691/231  
 GTG GCT GAC CTG GTA CAA GAG GTT CAG CAG GCC CGT GAC CTC CAG AAC AGG AGT GGG GCC  
 V A D L V Q E V Q Q A R D L Q N R S G A  
 721/241 751/251  
 ATG TCC CCG ATG TCA TGG AAC TCA GAC GCA TCT ACC TCC GAA GCG TCC TGA TGG GCC GCT  
 M S P M S W N S D A S T S E A S +  
 781/261 811/271  
 GCT TTG CCG TGG TGG ACC ACA GGC ATC TAC ACA GCC TGG ACT TTG GTT CTC TCC AGG AAG  
 841/281 871/291  
 GTA GCC CAG CAC TGT GAA GAC CCA GCA GGA AGC CAG GCT GAG TGA GCC ACA GAC CAC CTG  
 901/301 931/311  
 CTT CTG AAC TCA AGC TGC GTT TAT TAA TGC CTC TCC CCG ACC AGG CCG GCG TTG GCG CCT  
 961/321 991/331  
 GCA CAG ATA TTT CCA TTT CTT CCT CAC TAT GAC ACT GAG CAA GAT CTT GTC TCC ACT AAA  
 1021/341 1051/351  
 TGA GCT CCT GCG GGA GTA GTT GGA AAG TTG GAA CCG TGT CCA GCA CAG AAG GAA TCT GTG  
 1081/361 1111/371  
 CAG ATC AGC AGT CAC ACT GTT ACT CCA CAG CCG AGG AGA CCA GCT CAG AGG CCC AGG AAT  
 1141/381 1171/391  
 CCG AGC GAA GCA GAG AGG TGG AGA ACT GGG ATT TGA ACC CCC GCC ATC CTT CAC CAG AGC  
 1201/401 1231/411  
 CCA TGC TCA ACC ACT GTG CCG TTC TGC TGC CCC TGC AGT TGG CAG AAA GGA TGT TTT TGT  
 1261/421 1291/431  
 CCC ATT TCC TTG GAG GCC ACC GGG ACA GAC CTG GAC ACT AGG GTC AGG CCG GGT GCT GTG  
 1321/441 1351/451  
 GTG GGG AGA GGC ATG GCT GGG GTG GGG GTG GGG AGA CCT GGT TGG CCG TGG TCC AGC TCT  
 1381/461 1411/471  
 TGG CCC CTG TGT GAG TTG AGT CTC CTC TCT GAG ACT GCT AAG TAG GGG CAG TGA TGG TTG  
 1441/481 1471/491  
 CCA GGA CGA ATT GAG ATA ATA TCT GTG AGG TGC TGA TGA GTG ATT GAC ACA CAG CAC TCT  
 1501/501 1531/511  
 CTA AAT CTT CCT TGT GAG GAT TAT GGG TCC TGC AAT TCT ACA GTT TCT TAC TGT TTT GTA  
 1561/521 1591/531  
 TCA AAA TCA CTA TCT TTC TGA TAA CAG AAT TGC CAA GGC AGC GGG ATC TCG TAT CTT TAA  
 1621/541 1651/551  
 AAA GCA CTC CTC TTA TTC CTA AGG TAA TCC TAT TAA AAC ACA GCT TTA CAA CTT CCA TAT  
 1681/561  
 TAC AAA AAA AAA AAA AAA AAA

Figure 4

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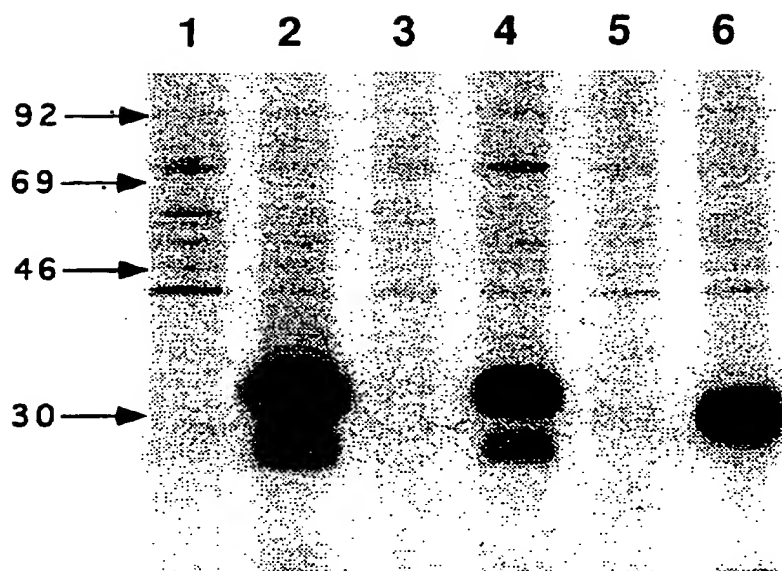


Figure 5

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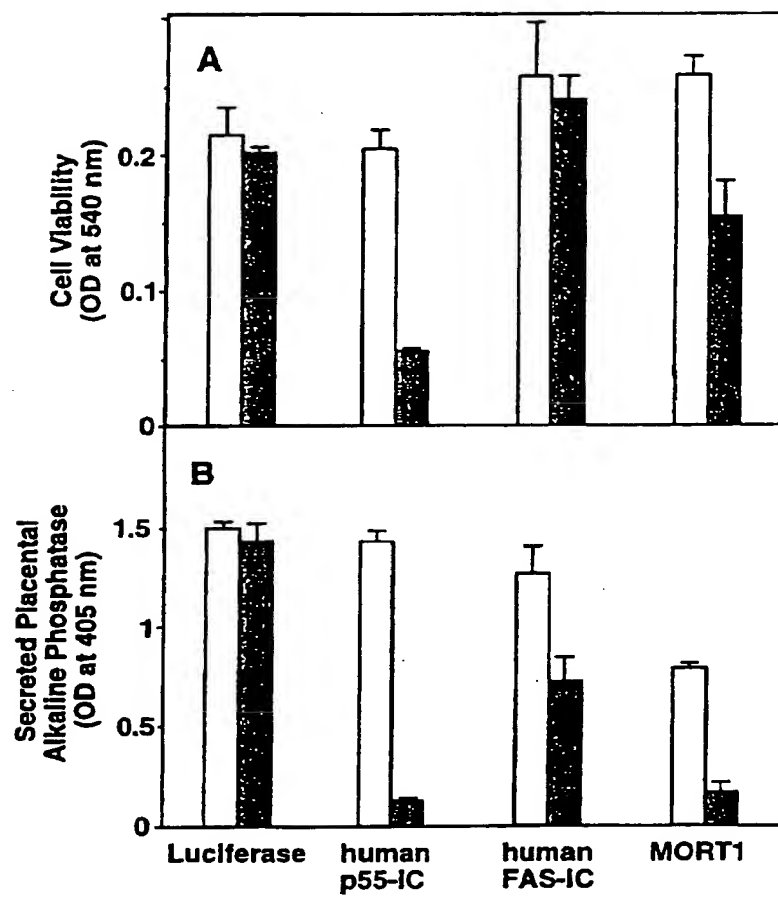


Figure 6

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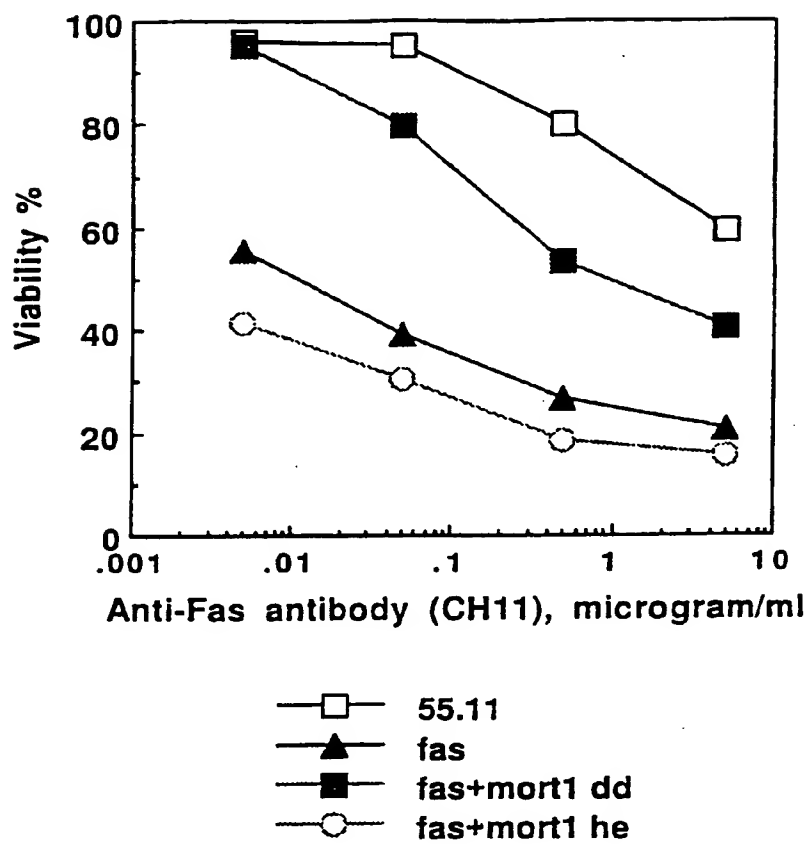


Figure 7